

Figure 1. mMRP amino acid sequence

1 MGSLFQEASP QAGTEQNKPT LASRFQQTILG DLLARLGSRG HYYVIHCLNP
51 TPGKIPGLLD VGHVAEQLRQ AGILEIIGTR STHFPVRVSF QVFLARFHAL
101 GSGRQKAASD QERCGAILSE VLGAESPLYH LGVTQVLLQE QGWQQLEQLW
151 AQRRSQALLT LHRGLRACIT RQRLRLLPRM QARVRGLQAR KRYLQRRSAL
201 GQLNTILLVA RPLLRRRQKL RCAPGPHSGE PWGKVSNDML GRLEIPAQLA
251 TLLERAEGHQ ALLTGSITES LPPEVPARPS LTLPPDIDQF PFSSEFVSTSF
301 QKPFLPRPGQ PLDEPLTRLD GENPQQALEI NRVMLRLLGE GSLQSWQEQT
351 MGTFLVQQAQ RRPGLRDEL F SQLVAQLWRN PDEQQNQRGW ALMVILLSSF
401 APTPALEKPL LKFVSDQAPS GMAALCQHKL LGALEQTPLA PMASRSHPTT
451 QLEWKAGLRR GRMALDVFTF NEESYSAEVE SWTTGEQFAG WILQSRGLEA
501 PPRGWSVSLH SGDAWRDLPG CDFVLDLIGQ TEDLGDPAGP HNYPTITPLGL
551 AESIPPAPGV QAPSLPPGLP PGPAPILASS RPPGEASKPE NLDGFVDHLF
601 EPALAPGFSD LEQGWALSRR MKGGGSGVGPT QQGYPMVYPG MVQAPSYQPA
651 MIPAPMPVMP AMGAVPTMPA MMVPPQPQPL VPSLDSRQLA LQQQNFINQQ
701 AMILAQQMTT QAMSLSLEQQ NQRHQHQAQT SGATSQPPPS TTAPKAKKPP
751 APQEKPEPNL EPSGVGLRED TPEEAESKPQ RPKSFQQKRD YFQKMGQDPI
801 RVKTKVKPPAK VQIPQEEMEE TEEEDETA E LSPPPPPPPV VKKPLKASRP
851 KAVKEDEAEP AQEEVPTQGE DPPVHSSNSA PQHKPKPSRPV PVQSSNSAPP
901 RPQPSREIRN IIRMYQSRPG PVAVPVQPTR PIKTFQKKND PKDEALAKLG
951 INGVHLPLST SPNQGKSSPP AVVPRPKARP RLEPSLSIQE KQGPLRLDLFG
1001 PCSNPPTAP APPPPPALPP PLSGEPKTPS VESHALTEPM EDKNISTKLL
1051 VPSGVCFSY ANAPWKFLR KEVFYPRENF SHPYCLSLLC QQILRDTTFE
1101 SCTRISQDER HKMKGLLDGL EVSLETLDIV EDSIKKRIVV AARDNWANYF

1151 SRIFPVSGES GSDVQLLGVS HRGLRLLKVT QSPSFHLDQL KTLCSYSYAE
1201 VLTVQCRGRS TLELSLKNEQ LILHTAWARA IKAMVDLFLS ELRKDSGYVI
1251 ALRSYITDDN SLLSFHRGDL IRLLPVTALE PGWQFGSAGG RSGLFPDDVV
1301 QPAAAPDLSF SLGKRNSWQR KSKLGPAQEV RKTEEVK*

1151 SRIFPVSGES GSDVQLLGVS HRGLRLLKVT QSPSFHLDQL KTLCSYSYAE

Figure 2. cDNA sequence of mMRP (variant 1)

1 CGCTGGGACT GTCACCTACC AGGTGCACAA GTTCATAAAC AGAAACAGGG
51 GCCACCTGGA CCCCGCTGTG CTGGAGATGC TCAGGCAGAG CCAGCTGCAG
101 GTGACCTAGC CTTCTTTTCA GTCATGGGC AGCCTGTTCC AAGAAGCAGA
151 GCCCCAGGCT GGGACTGAGC AAAACAAACC CACATTGGCC TCTCGATTCC
201 AGCAGACCCT GGGTGACTTG CTAGCTCGGC TAGGCAGCAG GGGCCATGTC
251 TACGTATCC ACTGTCTCAA TCCCACCCCT GGAAAGATCC CAGGCCTCTT
301 GGACGTGGGG CATGTGGCAG AGCAGCTGCG TCAGGCTGGC ATCTGGAGA
351 TCATAGGCAC CCGGAGTACC CACTTCCCCG TCGAGTGTC CTTCCAAGTC
401 TTTCTGGCAA GGTTCCATGC CCTGGGGTCA GGGAGACAGA AAGTCGCCTC
451 TGACCAGGAG AGGTGTGGTG CCATCCTCAG TGAAGTGCTG GGGCGCAGAT
501 CACCGCTGTA TCATCTTGGA GTCACCCAGG TCCTGCTGCA GGAACAGGGC
551 TGGCAGCAGC TAGAACAGCT GTGGGCTCAG CGGCCTCAC AGGCCCTGCT
601 CACTCTGCAC CGTGGCCTCC GAGCCTGTAT CACCCGGCAG CGCCTCCGTC
651 TCCTGCCCCG GATGCAGGCT CGTGTGCGTG GGCTCCAGGC CAGGAAGCGA
701 TATCTCCAGC GGAGGTCAGC TCTGGGACAG CTGAACACCA TTCTCCTAGT
751 GGCCCGGCC CTGCTCCGGA GACGACAGAA GCTACGGTGT GCCCTTGGCC
801 CGCACAGCGG GGAGCCCTGG GGGAAAGTGT CAAATATGGA CCTGGGTGCG
851 TTAGAGATCC CCGCCAGCT GGCTACTCTG CTGGAGAGGG CGGAAGGCCA
901 CCAGGCCTTG CTGACGGGGA GCATCACAGA GTCCCTGCCA CCTGAGGTCC
951 CCGCCCGGCC CAGCCTGACT CTCCTCCAG ACATTGACCA GTTTCCTTC
1001 TCCAGTTTTG TATCCACCAG CTTTCAGAAG CCATTCTGCT CTCGACCAGG
1051 GCAGCCACTG GACGAGCCCC TGACGCGGTT AGATGGCGAG AACCCCTCAGC

1101 AGGCTCTGGA GATCAACAGG GTGATGCTGC GGCTCCTGGG GGAAGGATCT
 1151 CTGCAGTCCT GGCAAGAGCA GACCATGGGC ACGTTCCTCG TGCAGCAGGC
 1201 CCAGCGACGG CCGGGACTCC GAGATGAGCT CTTCAGCCAG CTGGTGCCCC
 1251 AGCTGTGGCG CAACCCAGAT GAGCAACAGA ATCAGCGTGG CTGGGCCCTA
 1301 ATGGTGATCC TGCTCAGCTC CTTTGCTCCC ACACCTGCCC TGGAGAAGCC
 1351 ACTGCTCAAA TTTGTATCTG ACCAGGCTCC CAGTGGCATG GCAGCCCTGT
 1401 GCCAGCACAA GCTGTTAGGT GCCCTGGAGC AGACACCGCT GGCTCCCATG
 1451 GCTTCGAGGT CCCACCCACC CACACAACCT GAGTGAAGG CTGGTTTACG
 1501 TCGGGGCCCG ATGGCGCTGG ATGTGTTTAC ATTCAACGAG GAAAGTACT
 1551 CCGCGGAAGT GGAATCCTGG ACCACGGGAG AGCAGTTTGC AGGGTGGATC
 1601 CTACAGAGCA GAGGCCTGGA GCGCCCCCT CGTGGCTGGT CTGTGTCACT
 1651 GCATTCTGGG GATGCTTGGC GTGACTTGCC TGCTGTGAC TTTGTGTTGG
 1701 ACCTAATAGG CCAGACTGAG GACTTGGGAG ACCCAGCTGG TCCCCACAAC
 1751 TACCCCATCA CTCCTCTTGG TTTAGCTGAG AGCATCCCTC CAGCCCCCTG
 1801 TGTCCAGGCT CCTTCCCTGC CCCCAGGACT CCCTCCAGGT CCAGCCCCAA
 1851 TACTGGCCAG CAGCCGCCCT CCGGGCGAGG CCAGTAAGCC TGAGAACCTG
 1901 GATGTTTTCG TGGACCACCT CTTTGAACCA GCGCTCGCTC CGGGTTTCAG
 1951 TGATCTGGAA CAAGGCTGGG CCCTGAGCAG ACGCATGAAG GGAGGGGGGT
 2001 CTGTTGGGCC CACCCAGCAG GGCTACCCCA TGGTGATACC AGGTATGGTG
 2051 CAGGCACCTA GCTACCAGCC AGCTATGATA CCCGCACCGA TGCCCGTCAT
 2101 GCCAGCCATG GCGCGAGTCC CAACCATGCC AGCCATGATG GTGCCACCCC
 2151 AGCCACAGCC TCTGGTGCCC AGTTTGGACT CAAGGCAGCT GGCCTACAG
 2201 CAGCAAAACT TCATCAACCA GCAGGCGATG ATTCTGCGCG AGCAGATGAC
 2251 CACCCAGGCC ATGAGCCTGT CCCTGGAGCA GCAGAATCAG AGACACCAGC

2301 ACCAAGCTCA GACCTCTGGG GCCACCTCCC AGCCTCCACC CTCAACCACT
2351 GCTCCCAAGG CCAAGAAGCC TCCTGCCCCC CAAGAGAAGC CAGAGAGTAA
2401 CCTAGAGCCT TCGGGTGTG GCTTGAGAGA GGACACCCCA GAGGAAGCTG
2451 AAAGCAAGCC TCAGCGCCCC AAGAGCTTCC AACAGAAACG GGACTATTTC
2501 CAGAAGATGG GGCAAGATCC GATCAGAGTG AAGACGGTGA AACCTCCAGC
2551 CAAGGTTTCAG ATCCCCAAG AGGAGATGGA GGAGACGGAG GAGGAGGAGG
2601 ATGAGACCGC CGAGTTGTCC CCTCCTCCTC CCCCTCCCCC GGTGTGTAAG
2651 AAGCCGCTGA AGGCAAGCAG GCCCAAAGCC GTAAAGGAAG ATGAGGCAGA
2701 GCGCGCCGAG GAGGAAGTAC CGACCCAGGG CGAGGATCCC CCGGTGCACA
2751 GCTCCAACCT CGCACCTCAG CACCCCAAAC CCAGCAGGGT ACCCCCACTG
2801 CAGAGCTCCA ACTCCGCACC TCCACGCCCG CAACCCAGCA GGGAAATCCG
2851 AAACATCATC CGAATGTACC AGAGCCGTCC AGGGCCTGTG GCTGTGCCCCG
2901 TACAACCCAC CAGGCCCATC AAAAATTTTC AGAAGAAAAA TGACCTTAAG
2951 GATGAGGCTT TGGCTAAGTT AGGGATAAAT GCGCTCCACT TGCCCCCTATC
3001 GACATCGCCT AACCAAGGGA AGAGCTCTCC ACCGCTGTGA GTTCCTCGAC
3051 CTAAGGCTCG ACCTCGTCTT GAGCCTTCCC TATCCATCCA GGAAGAGCAG
3101 GGACCCCTTC GGGACTTGTG TGGCCCATGT AGTCCAAACC CACCTACAGC
3151 TCCAGCACCC CCGCCTCCAC CAGCACTCCC ACCGCTCTG TCTGGGGAGC
3201 CCAAGACCCC TTCAGTGGAG TCTCATGCCT TGACAGAGCC CATGGAGGAC
3251 AAGAACATCT CCACAAAGCT CTTTGTGCCC TCTGGAAGTG TGTGCTTCTC
3301 CTATGCCAAT GCACCTGGA AGTTGTCTT ACGCAAGGAG GTGTTCTACC
3351 CCCGGGAGAA CTTAGTCAT CCATACTGCC TCAGTCTCCT CTGCCAGCAG
3401 ATCCTGCGGG ACACCTTCAC AGAGTCCTGC ACCCGGATCT CACAGGATGA

3451	CGGGCACAAA	ATGAAAGGCC	TTCTGGGAGA	CTTGGAGGTG	AGTCTGGAGA
3501	CCCTTGACAT	TGTTGAAGAC	AGCATCAAAA	AACGCATCGT	GGTCGCTGCT
3551	CGGGACAAC	GGGCCAATTA	CTTCTCCCGC	ATCTTCCCAG	TCTCGGGTGA
3601	GAGTGGCAGC	GATGTACAGC	TGCTGGGTGT	GTCTCACC	GGACTGCGGC
3651	TGCTGAAGGT	GACCCAAAGC	CCGAGCTTCC	ACCTGGACCA	GCTGAAGACA
3701	CTCTGTTCCT	ACAGCTATGC	TGAAGTCTCT	ACCGTGCAGT	GCAGGGGCAG
3751	ATCCACCCTG	GAGCTGTCTT	TGAAGAATGA	GCAGCTGATA	CTGCACACAG
3801	CCTGGGCGAG	GGCCATCAAG	GCCATGGTGG	ATCTATTTCT	GAGTGAATCT
3851	AGGAAGGACT	CCGGCTATGT	CATCGCCCTG	CGCAGCTACA	TCACCGATGA
3901	CAATAGCCTC	CTCAGTTTCC	ACCGTGGGGA	CCTCATTAGG	TTACTGCCAG
3951	TGACCGCTCT	GGAACCAAGC	TGGCAGTTCG	GTTCTGCCCG	GGGCCGCTCC
4001	GGACTCTTTC	CCGATGACGT	GGTGCAGCCA	GCTGCTGCCC	CCGACCTCTC
4051	CTTTTCCCTG	GGAAAGAGAA	ACAGCTGGCA	ACGCAAGAGT	AAGCTGGGGC
4101	CAGCTCAGGA	GGTGAGGAAG	ACAGAAGAGG	TGAAGTGATA	CAGGCCTAAC
4151	TTGGAGACTG	AGAAGGAAAG	AGCAGGGTTG	CTTCGGGTGT	TGTCCACTTC
4201	CTGTCTGGT	GGCCAGGGCT	CAATGTGTTC	CTGTCCTTTA	CCATCTCCTG
4251	ACTTTTTGCC	ATTTGTGAGA	CTGTAAGTCA	CACCCTCTAA	CTCTGGTACT
4301	TAGTTCAGTG	TCTCCATAGA	GGATGCTTAA	TAAATAACCT	TGGTTTTCTT
4351	GGTTTCTGGT	GTCACTCCTC	TTGGGTCTAA	TGGGTATGGG	GACCAGGGCC
4401	TGAGAGTGAG	TATTGGGCCT	CTGGGCTAGA	TGGTGGGTAC	TGGGTGGGTA
4451	CCAAATTTC	TGTGCTCCCA	GCGCCCCACC	CATCCCAGGA	AACAAGAACC
4501	CAGTGAAGAC	TCGGAGGCCA	CCTCCTTTAC	AACCTACAGC	TCTTTGTCTG
4551	CCGACCCCCA	CAACTACACC	ATGCAGGAAT	TTGCCCTGCG	CTATTTCCGG
4601	AAGCCTCATA	CCTGGCTGAC	CCAGATGAGT	AGAGACACCA	AAGAGAAAGC

4651 TGCCATCAAC CTGATCCAGT AACTAAGGA CCCCATCCAG GAATCCCTTA
 4701 CCAGCTTCTG CAATGGGGAC ACAAACAGTA AAGCTGTGGC TGGCTTCAAG
 4751 GCTCTGATGC AGTTTATGGG GGACCAGCCT AAGCCCCGGG GCAAGGACGA
 4801 GCTGAGTCTG CTCTATGAGC TGCTGAAGCT GTGCCAAGAT GACCTTAGGG
 4851 ACGAGATGTA CTGCCAGGTC ATCAAGCAAG TCACAGGACA CCCCCAGCCA
 4901 AAGCACTGTG CTCTGGGCTG GAGCGTCCTC AGCCTCTTCA CAGGCTTCTT
 4951 TGCACCATCG ACCACGCTGA TGCCCTATGT GACCAAGTTC CTGCAGGATT
 5001 CCAGCCCCAG TGAAGAGTTG GCCAGGAGGA GCCAGGAGAA CCTCCAGCGC
 5051 ACAGTTAAAT ATGGGGGACG CCAGCAGCTG CCGTTACCTG GTGAAATGAA
 5101 TGCTTTTCTG AAAGGGCAAG CAGTTCGTTT GCTTCTAATT CACCTGCCTG
 5151 GGGGTGTGGA CTACAGGACG AATTCACAGA CATTCACAGT GGCAGGGGAA
 5201 GTGCTAGAGG AGCTGTGTGG ACAGATGGGC ATCACAGACT TGAAGAAGT
 5251 GCAGGAATTT GCCCTCTTTC TCATCAAAGG AGAAGGTGAG CTGGTTTCGGC
 5301 CGCTGTACCC CCATGAGTAC ATCAACAATG TGGTGACGGA CCAGGACATG
 5351 AGCCTTCACA GCCGACGGCT TGGTTGGGAG ACTCCACTGC ATTTTGATCA
 5401 CTCCACCTAC ACGGAAACCC ACTATGGCCA GGTGCTTCGG GACTACCTGC
 5451 AAGGGAAGCT GATAGTCAGC ACCCAGGCAG AGGCTCTACT TGCCAGCTT
 5501 GCTGCCTTCC AACACTTCGA CAAAACCGGA ACTTCTAGTC CTCCATCAGA
 5551 GCAAGAGCTG CTGTCTTATA TTCCCAAGCC ACTGCAATGG CAGGTGAACA
 5601 CAGCCAACAT AAAGAGCTTG GTGACCCAGG AGCTGAGGCA GATGCAAGGG
 5651 TACAGCAAGC AGAGAGCACA GATTGGCTTT ATAGAGAGCA CAGCGCAGCT
 5701 GCCCTCTTT GGCTACACTG TGTACGTAGT GCTGAGAGTG AGTAAGCTGG
 5751 CCCTCCCTGG ACCAGGCCTC CTGGGGCTGA ACCGTCAGCA CCTGGTCCTC

5801	ATGGACCCCA	GCTCTCAGGA	ACTCTGCTGC	TCTGTCATGC	TAAAGACCT
5851	GAAGCAGTTC	CACCTGTCTGA	GCCCACTGCA	GGAGGACGGG	CCCCCTGGCC
5901	TAGAACTCAA	CTATGGCTCT	GTTGACAACC	CCCAGACCAT	CTGGTTGGAG
5951	TTGCCACAGG	CCCAGGAGCT	GCAGCACACC	ATCATCTTCC	TGCTGGGCAG
6001	CATGTCCACT	CAGTGGCCAG	GTCTCTCTCTG	AGGAGTGGAG	ATAAGGCAGC
6051	GGTCTCTCAC	TGGGCAGTCT	GCCTTAGTCC	TGCTCTGAAT	CCGCTGCACA
6101	ACCCCCCACC	CCACGTGGAG	GCCAAAAGGC	AAAGTTGTGT	CACCTGGGAG
6151	AATAGGCAGA	CACATCCCTT	CTGGGGTGGGA	CTGCAACAGG	AGTTGGGGCA
6201	TTTGCTGGCT	AGCCCCAGGG	AAAATGCCCA	CCCAGCTCGA	AAGCGGCACA
6251	AGTAAACAC	CCAAGGAAAA	AAAAAAAAAA	AAAAAAAAAA	AAA

Figure 3. cDNA sequence of mMRP (variant 2)

1 CGCTGGGACT GTCACCTACC AGGTGCACAA GTTCATAAAC AGAAACAGGG
51 GCCACCTGGA CCCCGCTGTG CTGGAGATGC TCAGGCAGAG CCAGCTGCAG
101 GTGACCTAGC CTTCTTTCA GTCATGGGC AGCCTGTTCC AAGAAGCAGA
151 GCCCCAGGCT GGGACTGAGC AAAACAAACC CACATTGGCC TCTCGATTCC
201 AGCAGACCCT GGGTGACTTG CTAGCTCGGC TAGGCAGCAG GGGCCATGTC
251 TACGTATCC ACTGTCTCAA TCCCACCCCT GGAAAGATCC CAGGCCTCTT
301 GGACGTGGGG CATGTGGCAG AGCAGCTGCG TCAGGCTGGC ATCCTGGAGA
351 TCATAGGCAC CCGAGTACC CACTTCCCG TCGAGTGTG CTTCCAAGTC
401 TTTCTGGCAA GGTTCATGC CCTGGGTCA GGGAGACAGA AAGCTGCCTC
451 TGACCAGGAG AGGTGTGGTG CCATCCTCAG TGAAGTGTG GGGCAGAGT
501 CACCGCTGTA TCATCTTGA GTCACCCAGG TCCTGCTGCA GGAACAGGGC
551 TGGCAGCAGC TAGAACAGCT GTGGGCTCAG CGGCGCTCAC AGGCCCTGCT
601 CACTCTGCAC CGTGGCCTCC GAGCCTGTAT CACCCGGCAG CGCCTCCGTC
651 TCCTGCCCCG GATGCAGGCT CGTGTGCGTG GGCTCCAGGC CAGGAAGCGA
701 TATCTCCAGC GGAGGTCAGC TCTGGGACAG CTGAACACCA TTCTCCTAGT
751 GGCCCCGGCC CTGCTCCGGA GACGACAGAA GCTACGGTGT GCCCCTGGCC
801 CGCACAGCGG GGAGCCCTGG GGGAAAGTGT CAAATATGGA CCTGGGTCGC
851 TTAGAGATCC CCGCCAGCT GGCTACTCTG CTGGAGAGGG CGGAAGGCCA
901 CCAGGCCTTG CTGACGGGGA GCATCACAGA GTCCCTGCCA CCTGAGGTCC
951 CCGCCCGGCC CAGCCTGACT CTCCCTCCAG ACATTGACCA GTTCCCTTC
1001 TCCAGTTTTC TATCCACCAG CTTTCAGAAG CCATTCTGTC CTCGACCAGG
1051 GCAGCCACTG GACGAGCCCC TGACGCGGTT AGATGGCGAG AACCTCAGC

1101 AGGCTCTGGA GATCAACAGG GTGATGCTGC GGCTCCTGGG GGAAGGATCT
 1151 CTGCAGTCCT GGCAAGAGCA GACCATGGGC ACGTTCCTCG TGCAGCAGGC
 1201 CCAGCGACGG CCGGGACTCC GAGATGAGCT CTTCAGCCAG CTGGTGGCCC
 1251 AGCTGTGGCG CAACCCAGAT GAGCAACAGA ATCAGCGTGG CTGGGCCCTA
 1301 ATGGTGATCC TGCTCAGCTC CTTTGCTCCC ACACCTGCCC TGGAGAAGCC
 1351 ACTGCTCAAA TTTGTATCTG ACCAGGCTCC CAGTGGCATG GCAGCCCTGT
 1401 GCCAGCACAA GCTGTTAGGT GCCCTGGAGC AGACACCGCT GGCTCCCATG
 1451 GCTTCGAGGT CCCACCCACC CACACAACTT GAGTGGAAGG CTGGTTTACG
 1501 TCGGGGCCGC ATGGCGCTGG ATGTGTTTAC ATTCAACGAG GAAAGCTACT
 1551 CCGCGGAAGT GGAATCCTGG ACCACGGGAG AGCAGTTTGC AGGGTGGATC
 1601 CTACAGAGCA GAGGCCTGGA GGCGCCCCCT CGTGGCTGGT CTGTGTCACT
 1651 GCATTCTGGG GATGCTTGGC GTGACTTGCC TGGCTGTGAC TTTGTGTTGG
 1701 ACCTAATAGG CCAGACTGAG GACTTGGGAG ACCCAGCTGG TCCCCACAAC
 1751 TACCCCATCA CTCCTCTTGG TTTAGCTGAG AGCATCCCTC CAGCCCCTGG
 1801 TGTCCAGGCT CCTTCCCTGC CCCAGGACT CCCTCCAGGT CCAGCCCCAA
 1851 TACTGGCCAG CAGCCGCCCT CCGGGCGAGG CCAGTAAGCC TGAGAACCTG
 1901 GATGGTTTCG TGGACCACCT CTTTGAACCA GCGCTCGCTC CGGGTTTCAG
 1951 TGATCTGGAA CAAGGCTGGG CCCTGAGCAG ACGCATGAAG GGAGGGGGCT
 2001 CTGTTGGGCC CACCCAGCAG GGCTACCCCA TGGTGTAACC AGGTATGGTG
 2051 CAGGCACCTA GCTACCAGCC AGCTATGATA CCCGCACCGA TGCCCGTCAT
 2101 GCCAGCCATG GGCGCAGTCC CAACCATGCC AGCCATGATG GTGCCACCCC
 2151 AGCCACAGCC TCTGGTGCCC AGTTTGACT CAAGGCAGCT GGCACACTAG
 2201 CAGCAAAACT TCATCAACCA GCAGGCGATG ATTCTGGCGC AGCAGATGAC
 2251 CACCCAGGCC ATGAGCCTGT CCCTGGAGCA GCAGAATCAG AGACACCAGC

2301 ACCAAGCTCA GACCTCTGGG GCCACCTCCC AGCCTCCACC CTCAACCACT
 2351 GCTCCCAAGG CCAAGAAGCC TCCTGCCCCC CAAGAGAAGC CAGAGAGTAA
 2401 CCTAGAGCCT TCGGGTGTG GCTTGAGAGA GGACACCCCA GAGGAAGCTG
 2451 AAAGCAAGCC TCAGCGCCCC AAGAGCTTCC AACAGAAACG GGAATATTTT
 2501 CAGAAGATGG GGCAAGATCC GATCAGAGTG AAGACGGTGA AACCTCCAGC
 2551 CAAGGTTTCA ATCCCCAAG AGGAGATGGA GGAGACGGAG GAGGAGGAGG
 2601 ATGAGACCGC CGAGTTGTCC CCTCCTCCTC CCCCTCCCCC GGTGTGAAG
 2651 AAGCCGCTGA AGGCAAGCAG GCCCAAAGCC GTAAAGGAAG ATGAGGCAGA
 2701 GCCCGCCCAG GAGGAAGTAC CGACCCAGGG CGAGGATCCC CCGGTGCACA
 2751 GCTCCAATC CGCACCTCAG CACCCCAAAC CCAGCAGGGT ACCCCCAGTG
 2801 CAGAGCTCCA ACTCCGCACC TCCACGCCCG CAACCCAGCA GGGAAATCCG
 2851 AAACATCATC CGAATGTACC AGAGCCGTCC AGGGCCTGTG GCTGTGCCCC
 2901 TACAACCCAC CAGGCCCATC AAAACTTTTC AGAAGAAAAA TGACCCTAAG
 2951 GATGAGGCTT TGGCTAAGTT AGGGATAAAT GGCGTCCACT TGCCCTATC
 3001 GACATCGCCT AACCAAGGGA AGAGCTCTCC ACCGGCTGTA GTTCCTCGAC
 3051 CTAAGGCTCG ACCTCGTCTT GAGCCTTCCC TATCCATCCA GGAAAAGCAG
 3101 GGACCCCTTC GGGACTTGTT TGGCCCATGT AGTCCAAACC CACCTACAGC
 3151 TCCAGCACCC CCGCCTCCAC CAGCACTCCC ACCGCCTCTG TCTGGGAGC
 3201 CCAAGACCCC TTCAGTGGAG TCTCATGCCT TGACAGAGCC CATGGAGGAC
 3251 AAGAACATCT CCACAAAGCT CTTGTGCCC TCTGGAAGTG TGTGCTTCTC
 3301 CTATGCCAAT GCACCCTGGA AGTTGTCTT ACGCAAGGAG GTGTCTTACC
 3351 CCCGGGAGAA CTTAGTCAT CCATACTGCC TCACTCTCT CTGCCAGCAG
 3401 ATCCTGCGGG ACACCTTCAC AGAGTCCTGC ACCCGGATCT CACAGGATGA

3451 GCGGCACAAA ATGAAAGGCC TTCTGGGAGA CTTGGAGGTG AGTCTGGAGA
 3501 CCCTTGACAT TGTGTAAGAC AGCATCAAAA AACGCATCGT GGTGCTGCT
 3551 CGGACAACCT GGGCCAATTA CTTCTCCCGC ATCTTCCCAG TCTCGGGTGA
 3601 GAGTGGCAGC GATGTACAGC TGCTGGGTGT GTCTCACCAG GACTGCGGC
 3651 TGCTGAAGGT GACCCAAAGC CCGAGCTTCC ACCTGGACCA GCTGAAGACA
 3701 CTCTGTTTCT ACAGCTATGC TGAAGTCCTG ACCGTGCAGT GCAGGGGCG
 3751 ATCCACCCTG GAGCTGTCCT TGAAGAATGA GCAGCTGATA CTGCACACAG
 3801 CCTGGGCGAG GGCCATCAAG GCCATGGTGG ATCTATTCTT GAGTGAAGTC
 3851 AGGAAGGACT CCGGCTATGT CATCGCCCTG CGCAGCTACA TCACCGATGA
 3901 CAATAGCCTC CTCAGTTTCC ACCGTGGGGA CCTCATTAGG TTAGTGCCAG
 3951 TGACCCTCTT GGAACCAGGC TGGCAGTTCG GTTCTGCCGG GGGCCGCTCC
 4001 GGAATCTTTC CCGATGACGT GGTGCAGCCA GCTGCTGCCC CCGACCTCTC
 4051 CTTTTCCCTG GGAAAGAGAA ACAGCTGGCA ACGCAAGAGT AAGCTGGGGC
 4101 CAGCTCAGGA GGTGAGGAAG ACAGAAGAGG TGAAGTGATA CAGGCCTAAC
 4151 TTGGAGACTG AGAAGGAAAG AGCAGGGTTG CTTCGGGTGT TGTCCACTTC
 4201 CTGTCCTGGT GGCCAGGGCT CAATGTGTTT CTGTCCTTTA CCATCTCCTG
 4251 ACTTTTTCGC ATTTGTGAGA CTGTAAGTCA CACCCCTCTA CTCTGGTACT
 4301 TAGTTCAGTG TCTCCATAGA GGATGCTTAA TAAATAACCT TGGTTTTCTC
 4351 GGAAAAAAAA AAAAAAAAAA AAAAA

Figure 4. ORF HMRP1 partial amino acid sequence--longer clone
(437aa)

MYQSRPGPVFVPVQPSRPPKAFLRKIDPKDEALAKLINGAHSSPPMLSPSPGKGPPPAVAPRPKA
PLQLGPSSSIKEKQGPLDLDFGQKLPIAHTPPPPAPPLPLPEDPGTLSAERRCLTQPVEDQGVST
QLLAPSGSVCFSYTGTPWKLFLRKEVFYPRENFSPHYLRLLCEQILRDTFSESCIRISQNERRKM
KDLLGGLEVDLDSLTTTEDSVKKRIVVAARDNWANYFSRFFPVSGESGSDVQLLAVSHRGLRLLKV
TQGPGLRPDQLKILCSYSFAEVLGVECRGGSTLELSLKSEQLVLHTARARAIEALVELFLNELKKD
SGYVIALRSYITDNCSLLSFHRGDLIKLLPVATLEPGWQFGSAGGRSGLFPADIVQAAAAPDFSFS
KEQRSGWHKGQLSNGEPGLARWDRASEVRKMGEGQAEARPA

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Figure 5. hMRP1 partial DNA sequence--longer clone 4174 bp

CCGACAGCAGCAGGCTCGGGCCTCCGAGGCTGCGTCCCAGGCCCTCACCTCAGCCGTCACCTCCAAAG
CCCAGGAAGCCCCCACACCCCCGGAGAAGCCACAGCGTGACCTGGGATCAGAGGGTGGCTGCCTG
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GCTGAAGCTGTGCCAGCAGGAGAAGCTGAGGGATGAGATTACTGCCAGGTATCAAGCAGGTAC
AGGACACCCCCGCGCGAACACTGCACCTCGAGGCTGGAGCTTCTCAGCCTTCTCACAGGCTTCTT
CCCCCGTCGACCAGGCTGATGCCTACCTGACCAAGTTTCTGCAGGATTACAGGCCAGCCAAGA
GCTGGCCCCGAGCAGCCAGGAGCACCTCCAGCGCACAGTCAAATATGGGGGGCGCGGCGGATGCC
CCACCCGGGTGAAATGAAGGCTTTCCTGAAAGGACAAGCGATTTCGCTGCTTCTTATTACCTGCC
GGGGGTGTGGATTATAGGACGAATATCCAGACTTTCACAGTAGCAGCAGAAGTGCAGGAGGAGCT
GTGCCGGCAAATGGGTATCACGGAGCCTCAGGAAGTGCAGGAATTCGCCCTTCTTCCTCATCAAAGA
GAAGAGCCAGCTGGTGCGCCCCCTGCAGCCCCGCCAATACCTCAACAGCGTGGTAGTGGACCAGGA
CGTGAGCCTGCACAGCCGGCGGCTCCACTGGGAGACCCCACTGCACCTTCGATAACTCCACCTACAT
CAGCACCCACTACAGCCAGGTGCTGTGGGACTACCTTCAGGGGAAGCTGCCAGTCAGCGCCAAGGC
AGACGCGCAGCTCGCCAGGCTGGCCGCCCTGCAGCACCTCAGCAAGGCCAACAGGAATACCCCTC
AGGGCAGGACCTGCTAGCTTACGTGCCAAAGCAGCTGCAACGGCAGGTGAACACGGCTCCATCAA
GAACCTGATGGGTGAGGAGCTGAGACGGCTGGAAGGACACAGCCCCCAGGAAGCACAGATCAGCTT
CATTGAGGCCATGAGCCAGCTGCCCCCTCTTCGGCTACACCGTCTATGGGGTGTGCGAGTGAGCAT
GCAGGCCCTGTCCGACCCACTCTCTGGGGCTCAACCGCCAGCATCTCATCTCATGGACCCAG
CTCCAGAGCCTGTACTGCCGCATTGCCCTGAAGAGCCTGCAGCGCTCCACCTGCTAAGCCCTCT
GGAGGAGAAGGGGCCCCCTGGCCTGGAAGTCAACTATGGCTCAGTGACAACCCCCAGACCATCTG
GTTTGAGCTGCCACAGGCCAGGAGCTGCTATACACCACTGTCTTCTGATAGACAGCAGTGCCCTC
TTGCACTGAGTGGCCAGCATCAACTGAGAGGAGTGCAGGCCGGGAGAGAAGAGGATGAGGCTC
CCCCGCCCCAAGTCTACCCACATGGTCTGCCTTGGATGCTATCAGATCACTGTCTAGAACCTGC
CTCAGCACAGCCAGCCGGCCACATGCAGGCCATGAGGCAGGGGCTGCTATCAGTCAACAGCAG
GCAAAGAAAACAGCCAGACCCTCTCCAGGACGGCTGGGGCCAAAGCGGGCTGCAGGAACCTGGCT
GGGGCACCTGAGGTTGCCCAGTCTGAGGGAGATGCCACCCGACCCAGGCTCCGCCCCAGGCCCCA

CATTAGCACAAAGCCCAGGCATGGGAGAAACAGCTGCTGAGGAAATAAACTCCCTAAAAAAAAAAAA
AAAAAAAAAAAAAAAAA

[illegible]

Figure 6. ORF hMRP2 partial amino acid sequence --shorter clone
(786aa)

MYQSRPGPVFPVQPSRPPKAFLRKIDPKDEALAKLINGAHSSPPMLSPSPGKGPPPAVAPRPKA
PLQLGPSSSIKEKQGPLLDLFGQKLPIAHTPPPPAPPLPLPEDPGTLSAERRCLTQFVEDQGVST
QLLAPSGSVCFSTYGTGPWKLFRLKEVFYPRENFSPHYLRLLECEQILRDTFSESCIRISQNERRKM
KDLLGGLEVDLDSLTTTDESVKKRIVVAARDNWANYFSRFFPVSGESGSDVQLLAVSHRGLRLKKV
TQGPGLRPDQLKILCSYSFAEVLGVECRGGSTLELSLKSEQLVLHTARARAIEALVELFLNELKKD
SGYVIALRSYITDNCSSLSSFHRGDLIKLLPVATLEPGWQFGSAGGRSGLFPADIVQAAAAPDFSFS
KEQSGWHKQQLSNGEPLARWDRASERPAHPWSQAHSDDSEATSLSSVAYAFLPDSHSYTMQEFA
RRYFRRSQALLGQTDGGAAGKDTDSLQYTKAPIQESLLSLSDDVSKLAVASFLALMRFMGDQSKP
RGKDEMDLLYELLKLCQQEKLRLDEIYCQVIKQVTGHPREHCTRGWSFLSLLTGFFPSTRLMPYL
KFLQDSGPSQELARSSQEHQLQRTVKYGGRRRMPPPGEMKAFLKGQAIRLLLIHLPGGVDYRTNIQ
FTVAAEVQEELCRQMGITEPQEVQEFALFLIKEKSQVLRPLQPAEYLNSVVVDQDVSLSHGGSTG
RPHCTSITPPTSAPTARCCGTTFRGSCQSAQRTRSSPGWPPCSTSARPTGIPPQGRTC

Figure 7. HMRP2 partial DNA sequence--shorter clone (3780 bp)

CGGCAGCAGCAGGCTCGGGCCTCCGAGGCTGCGTCCCAGGCCCTCACCTCAGCCGTACCTCCAAG
CCCAGGAAGCCCCCACACCCCGGAGAACCCACAGCGTGACCTGGGATCAGAGGGTGGCTGCCTG
AGGGAGACCTCCGAGGAGGCTGAAGACAGGCCTATCAGCCCAAGAGCTTCCAGCAGAAACGGAAC
TATTTCCAGAGGATGGGGCAGCCACAGATCACAGTGAGGACGATGAAGCCCCCGGCCAAGGTCCAC
ATCCCCCAGGGGGAAGCGCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGCAGGAGGAGCAA
GAAGTGGAACAAGAGCAGCGCCGTCCCTCCTCCTCCCCCATCGTGAAGAAGCCATTGAAGCAA
GGTGGGGCCAAAGCTCCAAAAGAGGCTGAGGCTGAGCCAGCCAAGGAGACAGCGGCCAAGGGCCAT
GGCCAAGGGCCAGCCCAAGGCAGGGGGACTGTGGTGCGCAGTCAGACTCCAAGCCCAAGCGGCCAC
AACCAGCAGGGGAAATTGGCAACATCATCCGCATGTACCAGAGCCGCCCCGGCCCCCGTGCCTGTGC
CCGTGACGCCATCCAGGCCTCCCAAAGCTTTCTGAGGAAAATCGACCCCAAGGACGAGGCTCTGG
TCAAGCTGGGTATCAACGGTGCCCACTCGTCCCCGCCGATGCTGTCCCCCAGCCAGGAAAGGGCC
CCCCGCCAGCTGTGGCTCCTCGACCCAAGGCCCGCTACAGCTTGGGCCCTCTAGCTCCATCAAGG
AAAAGCAGGGGCCCTTCTGGACCTGTTTGGCCAGAAGCTGCCTATTGCCACACACCCCCACCTC
CACCAGCGCCACCACTGCCTCTGCCCCGAGGACCCAGGGACCCCTTTCAGCAGAGCGTCGTGTGTTGA
CAGAGCCGTGGAGGACCAGGGGTCTCCACCCAGCTACTCGCGCCCTCTGGCAGCGTGTGCTTCT
CCTACACCGGCACGCCCTGGAAGTTGTTCTTACGCAAGGAGGTGTTCTACCCACGGGAGAACTTCA
GCCATCCCTACTACCTGAGGCTCCTCTGTGAGCAGATCCTACGGGACACCTTCTCCGAGTCTGTGA
TCCGGATTTCACAGAATGAGCGCGGAAAATGAAAGACCTGCTGGGAGGCTTGGAGGTGGACCTGG
ATTCTCTCACCACCACGAAGACAGCGTCAAGAAGCGCATCGTGGTGGCCGCTCGGGACAAGTGGG
CAATTACTTCTCCCGCTTCTTTCTGTCTCGGGGAGAGTGCGACGACGTGCAGCTGTTAGCCG
TGTCCCACCGTGGGCTGCGACTGCTCAAGGTGACCCAAGGCCCGGCCCTCCGCCCGACCAGCTGA
AGATTCTCTGCTCATACAGCTTTGCGGAGGTGCTGGGTGTGGAGTGCCGGGGCGGCTCCACCTGG
AGCTGTCACTGAAGAGCGAGCAGCTGGTGTGTCACACAGCCCGGGCAAGGGCCATCAGGGCGTGG
TTGAGTATTCTGAATGAGCTTAAGAAGGACTCCGGCTATGTATCGCCCTGCGCAGCTACATCA
CTGACAACTGCAGCTCCTCAGCTTCCACCGTGGGGACCTCATCAAGTGTGCTGCCGTGGCCACCC
TGGAGCCAGGCTGGCAGTTTGGCTCTGCGGGGGCCGTTCCGGACTCTTTCTGCGACATAGTGC
AGCGGCTGCCGCTCCCGACTTTTCTTCTCCAAGGAGCAGAGGAGTGGCTGGCACAAGGTCAGC
TGTCCAACGGGGAACAGGGCTGGCTCGGTGGGACAGGGCCTCAGAGCGCCTGCCACCCCTTGGGA
GCCAGGCACACAGTGACGACTCGAGGCCACCAGCCTGTCTCTGTGGCCTATGCCTTTCTGCCCG
ACTCCACAGCTACACCATGCAAGGAATTCGCCCCGCGTTACTTCCGGAGGTCCAGGCCCTGTCTGG
GCCAGACTGATGGAGGTGCCGAGGAAGGACACGGACAGCCTGGTGCAGTACCAAGGCTCCCA

TCCAGGAGTCGCTCCTCAGCCTCAGTGATGATGTGAGCAAGCTGGCTGTAGCCAGCTTCCTGGCCC
TGATGCGGTTTATGGGTGACCACTCCAAGCCCCGGGGCAAGGATGAGATGGATCTGCTCTATGAAC
TGCTGAAGCTGTGCCAGCAGGAGAAGCTGAGGGATGAGATTTACTGCCAGTTATCAAGCAGGTCA
CAGGACACCCCCGGCCGGAACACTGCACCTCGAGGCTGGAGCTTCCTCAGCCTTCTCACAGGCTTCT
TCCCCCGCTGCACCAGGCTGATGCCCTACCTGACCAAGTTTCTGCAAGATTTCAGGCCCCAGCCAAG
AGCTGGCCCCGAGCAGCCAGGAGCACCTCCAGCGCACAGTCAAATATGGGGGGCGCCGGCGGATGC
CCCCACCGGGTGAAATGAAGGCTTTCCTGAAAGGACAAGCGATTTCGCTGCTTCTTATTCACTGC
CGGGGGGTGTGGATTATAGGACGAATATCCAGACTTTCACAGTAGCAGCAGAAGTGCAGGAGGAGC
TGTGCCGCAAAATGGGTATCACGGAGCCTCAGGAAGTGCAGGAATTGCCCCCTTCTCCTCATCAAAG
AGAAGAGCCAGCTGGTGGGCCCTCAGCCCCGCCGAATACCTCAACAGCTGGTAGTGGACCAGG
ACGTGAGCCTGCACAGCGCGGCTCCACTGGGAGACCCACTGCACCTTCGATAACTCCACCTACAT
CAGCACCCACTACAGCCAGGTGCTGTGGGACTACCTTCAGGGGAAGCTGCCAGTCAGCGCCAAGGC
AGACGCGCAGCTCGCCAGGCTGGCCGCCCTGCAGCACCTCAGCAAGGCCAACAGGAATACCCCTC
AGGGCAGGACCTGTAGCTTACGTGCCAAAGCAGCTGCAACGGCAGGTGAACACGGCCTCCATCAA
GAACCTGATGGGTGAGGAGCTGAGACGGCTGGAAGGACACAGCCCCAGGAAGCACAGATCAGCTT
CATTGAGGCCATGAGCCAGCTGCCCTCTTCGGCTACACCGTCTATGGGGTCTGCGAGTGAGCAT
GCAGGCCCTGTCCGGACCCACTCTCCTGGGGCTCAACCGCCAGCATCTCATCTCATGGACCCAG
CTCCAGAGCCTGTACTGCCGATTGCCCTGAAGAGCCTGCAGCGGCTCCACCTGCTAAGCCCTCT
GGAGGAGAAGGGGCCCCCTGGCCTGGAAGTCAACTATGGCTCAGCTGACAACCCCCAGACCCTCG
GTTTGAGCTGCCACAGGCCAGGAGCTGCTATACCACTGTCTTCCTGATAGACAGCAGTGCCCTC
TTGCACTGAGTGGCCAGCATCAACTGAGAGGAGTGCAGGCCGGGGAGAGAAGAGGATGAGGCCTC
CCCCGGCCCAAGTCTCACCCACATGGTCTGCCTTGGATGCTATCAGATCACTGTCTAGAACCTGC
CTCAGCACAGCCAGCGGCCACATGCAGGCCATGAGGCAGGGGCTGCTATCAGCTCACCAGCAG
GCAAGAAAAACAGCCAGACCCTCTCCAGGACGGCCTGGGGCCAAGCGGGCTGCAGGAACCTCGGCT
GGGGCACCTGAGGTTGCCAGTCTGAGGGAGATGCCACCCGACCCAGGCTCCGCCCAGGCCCCA
CATTAGCACAAAGCCAGGCATGGGAGAAACAGCTGCTGAGGAAATAAACTCCCTAAAAAAAAAA
AAAAAAAAAAAAAAAAAAAA

Figure 8.

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mMRP: 914  MYQSRPGPVAVPVQPTRIKTFQKKNDPKDEALAKLINGVHL-PLSTSPNQKGKSSPPAV 972
MYQSRPGPV VPVQP+RP K F +K DPKDEALAKLING H P SP+ GK PPAV
hMRP: 1    MYQSRPGFPVPVPQSPRPKAFLRKIDPKDEALAKLINGAHSSPMLSPSPGKGPPPAV 60

mMRP: 973  VPRPKARPRLEPSLSIQEKQGPLRDLFGPCSPNPPTAPAPPPPPALPPLSGEPKTPSVE 1032
PRPKA +L PS SI+EKQGPL DLFQ P A PPPPPA P PL +P T S E
hMRP: 61   APRPKAPLQLGPPSSSIKEKQGPLEDLFGQ---KLPIAHTPPPPAPPLPLPEDPGTLSAE 117

mMRP: 1033 SHALTEPMEDKNISTKLLVPSGSVCFSYANAPWKFLRKEVFYPRENFSHFYCLSLCQQ 1092
LT+P+ED+ +ST+LL PSGSVCFSY PWKFLRKEVFYPRENFSHFY L LLC+Q
hMRP: 118  RRCLTPQVEDQGVSTQLLAPSGSVCFSYTGTPWKFLRKEVFYPRENFSHFYLRLLCEQ 177

mMRP: 1093 ILRDTFTESCTRISQDERHKMKGLLGDLEVSLETLDIVEDSIKKRIVVAARDNWANYFSR 1152
ILRDTF+ESC RISQ+ER KMK LLG LEV L++L EDS+KKRIVVAARDNWANYFSR
hMRP: 178  ILRDTFESCTRISQNERRKMKDLLGGLVDLDSLTSTEDSVKKRIVVAARDNWANYFSR 237

mMRP: 1153 IFPVSGESGSDVQLLGVSHRGLRLLKVTQSPSFHLDQLKTLCSYSYAEVLTVQCRGRSTL 1212
FPVSGESGSDVQLL VSHRGLRLLKVTQ P DQLK LCSYS+AEVL V+CRG STL
hMRP: 238  FFPVSGESGSDVQLLAVSHRGLRLLKVTQSGPLRPDQLKILCSYSFAEVLGVECRGGSTL 297

mMRP: 1213 ELSLKNQELILHTAWARAIAKAMVDLFLSELKDKSGYVIALRSYITDDNSLLSFHRGDLIR 1272
ELSLK+EQ+LHTA ARAI+A+V+LFL+EL+KDSGYVIALRSYITD+ SLLSFHRGDLI+
hMRP: 298  ELSLKSEQVLHTARARAIEALVELFLNELKKDSCGYVIALRSYITDNCSLLSFHRGDLIK 357

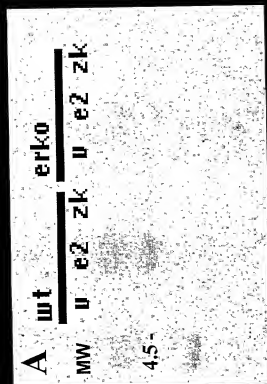
mMRP: 1273 LLPVTALEPGWQFGSAGGRSGLFPDDVVQPAAPDLSFSLGRNSWQR 1320
LLPV LEPGWQFGSAGGRSGLFP D+VQPAAPD SFS +R+ W +
hMRP: 358  LLPVATLEPGWQFGSAGGRSGLFPADIVQPAAPDFSFSKEQRSGWHK 405

```

Identities = 302/408 (74%), Positives = 334/408 (81%), Gaps = 4/408 (0%)



FIGURE 9



Regulation of mMRP Genes by Estrogen.

Northern hybridization of liver RNA from WT and ERKO mice treated with vehicle (V), 17β-estradiol (E2), and antiestrogen ZK compound. The myosin-related protein gene was only detected after E2 treatment in WT mouse.

FIGURE 10



Tissue Specific Regulation of mMRP by Estrogen.
 RT-PCR was performed on total RNA from WT or ERKO liver and brain tissues treated with vehicle (V), 17 β -estradiol (E2), and ZK compound. RNA quantity was controlled by RT-PCR on a house-keeping gene (GAPDH) in the same experiment.

FIGURE 11

Chromosomal Localization of Mouse MRP

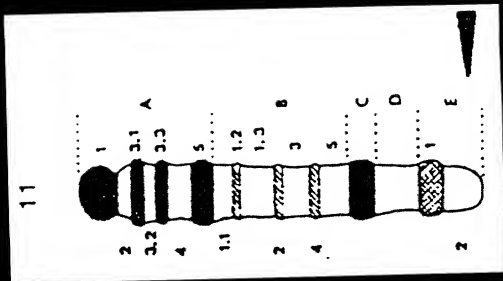


FIGURE 12

Chromosomal Localization of Human MRP

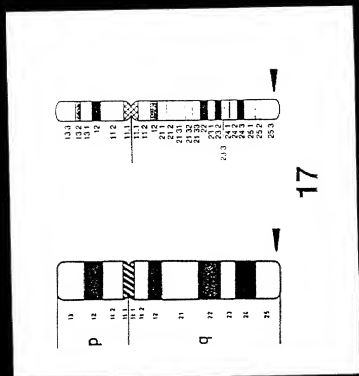


FIGURE 13